

SEQUENCE LISTING <110> Mack, Da Gish, Kurt EOS Biotechnology, Inc. <120> Methods of Diagnosis of Breast Cancer, Compositions and Methods of Screening for Modulators of Breast Cancer <130> 018501-001200US <140> US 09/829,472 <141> 2001-04-09 <150> US 09/525,361 <151> 2000-03-15 <160> 20 <170> PatentIn Ver. 2.1 <210> 1 <211> 3213 <212> DNA <213> Homo sapiens <220> <223> BCA4, osteoblast specific factor 2 (periostin) <220> <221> CDS <222> (12)..(2522) <223> BCA4 <400> 1 agagactcaa gatgattccc tttttaccca tgttttctct actattgctg cttattgtta 60 accctataaa cgccaacaat cattatgaca agatcttggc tcatagtcgt atcaggggtc 120 gggaccaagg cccaaatgtc tgtgcccttc aacagatttt gggcaccaaa aagaaatact 180 tcagcacttg taagaactgg tataaaaagt ccatctgtgg acagaaaacg actgttttat 240 atgaatgttg ccctggttat atgagaatgg aaggaatgaa aggctgccca gcagttttgc 300 ccattgacca tgtttatggc actctgggca tcgtgggagc caccacaacg cagcgctatt 360 ctgacgcctc aaaactgagg gaggagatcg agggaaaggg atccttcact tactttgcac 420 cgagtaatga ggcttgggac aacttggatt ctgatatccg tagaggtttg gagagcaacg 480 tgaatgttga attactgaat gctttacata gtcacatgat taataagaga atgttgacca 540 aggacttaaa aaatggcatg attattcctt caatgtataa caatttgggg cttttcatta 600 accattatcc taatggggtt gtcactgtta attgtgctcg aatcatccat gggaaccaga 660 ttgcaacaaa tggtgttgtc catgtcattg accgtgtgct tacacaaatt ggtacctcaa 720 ttcaaqactt cattqaaqca qaagatgacc tttcatcttt tagagcagct gccatcacat 780 cggacatatt ggaggccctt ggaagagacg gtcacttcac actctttgct cccaccaatg 840 aggettttga gaaaetteea egaggtgtee tagaaaggtt catgggagae aaagtggett 900 ccgaagctct tatgaagtac cacatcttaa atactctcca gtgttctgag tctattatgg 960

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Tyr His Arg Arg Lys Ala Gln Gln Met Thr Gln Lys Tyr Glu Glu Glu Leu Thr Leu Thr Arg Glu Asn Ser Ile Arg Arg Leu His Ser His His 395 390 Ser Asp Pro Arg Ser Gln Pro Glu Glu Ser Val Gly Leu Arg Ala Glu 405 410 Gly His Pro Asp Ser Leu Lys Asp Asn Ser Ser Cys Ser Val Met Ser 425 Glu Glu Pro Glu Gly Arg Ser Tyr Ser Thr Leu Thr Thr Val Arg Glu 440 Ile Glu Thr Gln Thr Glu Leu Leu Ser Pro Gly Ser Gly Arg Thr Glu 455 Glu Asp Asp Asp Gln Asp Glu Gly Ile Lys Gln Ala Met Asn His Leu 475 Cys Arg Lys Met Gly Pro 485 <210> 8 <211> 3085 <212> DNA <213> Homo sapiens <220> <223> BCZ6, IL-6 receptor beta chain (gp130, oncostatin M, IL-11, LIF and CNTF receptor) type I transmembrane protein <220> <221> CDS <222> (256)..(3012) <223> BCZ6 <400> 8 gagcagccaa aaggcccgcg gagtcgcgct gggccgcccc ggcgcagctg aaccgggggc 60 cgcgcctgcc aggccgacgg gtctggccca gcctggcgcc aaggggttcg tgcgctgtgg 120 agacgcggag ggtcgaggcg gcgcggcctg agtgaaaccc aatggaaaaa gcatgacatt 180 tagaagtaga agacttagct tcaaatccct actccttcac ttactaattt tgtgatttgg 240 aaatatccgc gcaagatgtt gacgttgcag acttgggtag tgcaagcctt gtttattttc 300 ctcaccactg aatctacagg tgaacttcta gatccatgtg gttatatcag tcctgaatct 360 ccagttgtac aacttcattc taatttcact gcagtttgtg tgctaaagga aaaatgtatg 420 gattattttc atgtaaatgc taattacatt gtctggaaaa caaaccattt tactattcct 480 aaggagcaat atactatcat aaacagaaca gcatccagtg tcacctttac agatatagct 540 tcattaaata ttcagctcac ttgcaacatt cttacattcg gacagcttga acagaatgtt 600 tatggaatca caataatttc aggcttgcct ccagaaaaac ctaaaaattt gagttgcatt 660 gtgaacgagg ggaagaaaat gaggtgtgag tgggatggtg gaagggaaac acacttggag 720 acaaacttca ctttaaaatc tgaatgggca acacacaagt ttgctgattg caaagcaaaa 780 cgtgacaccc ccacctcatg cactgttgat tattctactg tgtattttgt caacattgaa 840 gtctgggtag aagcagagaa tgcccttggg aaggttacat cagatcatat caattttgat 900 cctgtatata aagtgaagcc caatccgcca cataatttat cagtgatcaa ctcagaggaa 960 ctgtctagta tcttaaaatt gacatggacc aacccaagta ttaagagtgt tataatacta 1020 aaatataaca ttcaatatag gaccaaagat gcctcaactt ggagccagat tcctcctgaa 1080 gacacagcat ccacccgatc ttcattcact gtccaagacc ttaaaccttt tacagaatat 1140 gtgtttagga ttcgctgtat gaaggaagat ggtaagggat actggagtga ctggagtgaa 1200

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Thr Thr Glu Ser Thr Gly Glu Leu Leu Asp Pro Cys Gly Tyr Ile Ser
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                                                      30
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395

390

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Thr Glu Gly His Ser Ser Gly Ile Gly Gly Ser Ser Cys Met Ser Ser 725 Ser Arg Pro Ser Ile Ser Ser Ser Asp Glu Asn Glu Ser Ser Gln Asn Thr Ser Ser Thr Val Gln Tyr Ser Thr Val Val His Ser Gly Tyr Arg 760 His Gln Val Pro Ser Val Gln Val Phe Ser Arg Ser Glu Ser Thr Gln Pro Leu Leu Asp Ser Glu Glu Arg Pro Glu Asp Leu Gln Leu Val Asp 785 His Val Asp Gly Gly Asp Gly Ile Leu Pro Arg Gln Gln Tyr Phe Lys 810 805 Gln Asn Cys Ser Gln His Glu Ser Ser Pro Asp Ile Ser His Phe Glu Arg Ser Lys Gln Val Ser Ser Val Asn Glu Glu Asp Phe Val Arg Leu 840 Lys Gln Gln Ile Ser Asp His Ile Ser Gln Ser Cys Gly Ser Gly Gln 855 Met Lys Met Phe Gln Glu Val Ser Ala Ala Asp Ala Phe Gly Pro Gly 875 Thr Glu Gly Gln Val Glu Arg Phe Glu Thr Val Gly Met Glu Ala Ala Thr Asp Glu Gly Met Pro Lys Ser Tyr Leu Pro Gln Thr Val Arg Gln 905 Gly Gly Tyr Met Pro Gln 915 <210> 10 <211> 4215 <212> DNA <213> Homo sapiens <223> BFG4, KIAA0882 protein type II membrane protein <220> <221> CDS <222> (108)..(2777) <223> BFG4 <400> 10 gaacttatgt agcctcatta tecegeteeg tgaggtgaca attgtggaaa aggcagacag 60 ctccagtgtg ctccccagtc ccttatcaca tcagcacccg aaacaggatg accttcctat 120 ttqccaactt gaaagataga gactttctag tgcagaggat ctcagatttc ctgcaacaga 180 ctacttccaa aatatattct gacaaggagt ttgcaggaag ttacaacagt tcagatgatg 240 aggtgtactc tcgacccagc agcctcgtct cctccagccc ccagagaagc acgagctctg 300 atgctgatgg agagcgccag tttaacctaa atggcaacag cgtccccaca gccacacaga 360 ccctgatgac catgtatcgg cggcggtctc ccgaggagtt caacccgaaa ttggccaaag 420

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<400> 18

Leu Arg Leu Lys Cys Arg Asn Ala Cys Cys Gln Arg Trp Tyr Phe Thr

Phe Asn Gly Ala Glu Cys Ser Gly Pro Leu Pro Ile Glu Ala Ile Ile 165 170 175

Tyr Leu Asp Gln Gly Ser Pro Glu Met Asn Ser Thr Ile Asn Ile His 180 185 190

Arg Thr Ser Ser Val Glu Gly Leu Cys Glu Gly Ile Gly Ala Gly Leu 195 200 205

Val Asp Val Ala Ile Trp Val Gly Thr Cys Ser Asp Tyr Pro Lys Gly 210 215 220

Asp Ala Ser Thr Gly Trp Asn Ser Val Ser Arg Ile Ile Ile Glu Glu 225 230 235 240

Leu Pro Lys

<210> 20

<211> 243

<212> PRT

<213> Mus sp.

<220>

<223> mouse BCN4, ESTs, mouse orthologue of human BCN4

<220>

<221> MOD\_RES

<222> (1)..(243)

<223> Xaa = any amino acid

<400> 20

Xaa Xaa Xaa Ala Ala Pro Pro Gln Leu Leu Gly Leu Phe Leu 1 5 10 15

Val Leu Leu Leu Leu Gln Leu Ser Ala Pro Ser Ser Ala Ser Glu 20 25 30

Asn Pro Lys Val Lys Gln Lys Ala Leu Ile Arg Gln Arg Glu Val Val 35 40 45

Asp Leu Tyr Asn Gly Met Cys Leu Gln Gly Pro Ala Gly Val Pro Gly 50 60

Arg Asp Gly Ser Pro Gly Ala Asn Gly Ile Pro Gly Thr Pro Gly Ile 65 70 75 80

Pro Cys Gln Asp Gly Phe Lys Gly Glu Lys Gly Glu Cys Leu Arg Glu 85 90 95

Ser Phe Glu Glu Ser Trp Thr Pro Asn Tyr Lys Gln Cys Ser Trp Ser 100 105 110

Ser Leu Asn Tyr Gly Ile Asp Leu Gly Lys Ile Ala Glu Cys Thr Phe 115 120 125

Thr Lys Met Arg Ser Asn Ser Ala Leu Arg Val Leu Phe Ser Gly Ser 130 135 140

Leu Arg Leu Lys Cys Arg Asn Ala Cys Cys Gln Arg Trp Tyr Phe Thr 145 150 155 160

Phe Asn Gly Ala Glu Cys Ser Gly Pro Pro Pro Ile Glu Ala Ile Xaa 165 170 175

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Ser Asp Tyr Pro Lys Gly 210 215 220

Asp Ala Tyr Thr Gly Trp Asp Ser Val Ser Arg Ile Ile Ile Glu Glu 225 230 235 240

Leu Pro Lys